

SEQUENCE LISTING

<110> Ben-Sasson, Shmuel A.

<120> Short Peptides Which Selectively
Modulate the Activity of Protein Kinases

<130> 1242.1029-000 (CMCC-679)

<140> US 09/161,094

<141> 1998-09-25

<160> 172

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Gly Asn Pro Gly Phe
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 Lys Ala Val

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<400> 27
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 1 5 10 15
 Lys Ser Ile

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 Lys Val Leu

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 Lys Val Leu

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 Val Lys Pro Phe
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 Glu Pro Gly Val
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 1 5 10 15
 Thr Gly Lys Tyr Leu
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1 5 10 15
Asp Gly Lys Tyr Leu
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1 5 10 15
Glu Gly Arg Ala Leu
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Glu Gly Gln Asp Leu
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Glu Gly Gly Lys Val
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Thr Glu Phe Met Ala Lys Gly Ser Leu Leu Asp Phe Leu Lys Ser Asp
 1 5 10 15
 Glu Gly Ser Lys Gln
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<210> 44

<211> 21

<212> PRT

<213> unknown

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Thr Glu Tyr Met Glu Asn Gly Ser Leu Val Asp Phe Leu Lys Thr Pro
 1 5 10 15
 Ser Gly Ile Lys Leu
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Thr Glu Tyr Met Ala Lys Gly Ser Leu Val Asp Tyr Leu Arg Ser Arg
 1 5 10 15
 Gly Arg Ser Val Leu
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 1 5 10 15
 Gly Arg Ala Leu Val
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 Met Glu Leu Cys Thr Leu Gly Glu Leu Arg Ser Phe Leu Gln Val Arg
 1 5 10 15
 Lys Tyr Ser Leu
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 1 5 10 15
 Asn Arg Gln Glu Val
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 Ile Glu Tyr Ala Pro Tyr Gly Asn Leu Leu Asp Phe Leu Arg Lys Ser
 1 5 10 15
 Arg Val Leu Glu Thr Asp Pro Ala Phe Ala Arg Glu His Gly Thr Ala
 20 25 30
 Ser Thr Leu
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<400> 50
 Ile Glu Tyr Ala Pro His Gly Asn Leu Leu Asp Phe Leu Arg Lys Ser
 1 5 10 15

Arg Val Leu Glu Thr Asp Pro Ala Phe Ala Ile Ala Asn Ser Thr Ala
 20 25 30
 Ser Thr Leu
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 1 5 10 15
 Arg Pro Pro Gly Leu Glu Tyr Cys Tyr Asn Pro Ser His Asn Pro Glu
 20 25 30
 Glu Gln Leu
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<400> 52
 Val Glu Tyr Ala Ser Lys Gly Asn Leu Arg Glu Tyr Leu Arg Ala Arg
 1 5 10 15
 Arg Pro Pro Gly Met Glu Tyr Ser Tyr Asp Ile Asn Arg Val Pro Glu
 20 25 30
 Glu Gln Met
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 20 25 30
 Glu Gln Leu
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 Arg Pro Pro Gly Pro Asp Leu Ser Pro Asp Gly Pro Arg Ser Ser Glu
 20 25 30
 Gly Pro Leu
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 Thr Glu Tyr Cys Phe Tyr Gly Asp Leu Val Asn Tyr Leu His Lys Asn
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 Arg Asp Ser Phe Leu Ser His His Pro Glu Lys Pro Lys Lys Glu Leu
 20 25 30
 Asp Ile Phe Gly Leu Asn Pro Ala
 35 40

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 Thr Glu Tyr Cys Arg Tyr Gly Asp Leu Val Asp Tyr Leu His Arg Asn
 1 5 10 15
 Lys His Thr Phe Leu Gln His His Ser Asp Lys Arg Arg Pro Pro Ser
 20 25 30
 Ala Glu Leu Tyr Ser Asn Ala Leu
 35 40

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 1 5 10 15
 Glu Arg Ser Pro
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<210> 62
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<400> 62
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 1 5 10 15
 Gln Arg Asn Pro
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<210> 63
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<400> 63
 Thr Gln Leu Met Pro Phe Gly Cys Leu Leu Asp Tyr Val Arg Glu His
 1 5 10 15
 Lys Asp Asn Ile
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<400> 64
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 1 5 10 15
 Arg Gly Arg Leu

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 Thr Gln Tyr Leu Pro Leu Gly Ser Leu Leu Asp His Val Arg Gln His
 1 5 10 15
 Arg Gly Ala Leu
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ERB3
 ERB4
 ERBB2
 ERBB3
 ERBB4
 ERBB5
 ERBB6
 ERBB7

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<400> 66
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 1 5 10 15
 Lys Asp Asn Ile
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<210> 67
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<400> 67
 Val Glu Tyr Ala Lys Tyr Gly Ser Leu Arg Gly Phe Leu Arg Glu Ser
 1 5 10 15
 Arg Lys Val Gly Pro Gly Tyr Leu Gly Ser Gly Ser Arg Asn Ser
 20 25 30
 Ser Ser Leu Asp His Pro Asp Glu Arg Ala Leu
 35 40

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 Phe Glu Tyr Met Arg His Gly Asp Leu Asn Arg Phe Leu Arg Ser His
 1 5 10 15
 Gly Pro Asp Ala Lys Leu Leu Ala Gly Gly Glu Asp Val Ala Pro Gly
 20 25 30
 Pro Leu

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<220>
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<400> 69
 Phe Glu Tyr Met Lys His Gly Asp Leu Asn Lys Phe Leu Arg Ala His
 1 5 10 15
 Gly Pro Asp Ala Val Leu Met Ala Glu Gly Asn Pro Pro Thr Glu Leu
 20 25 30

<210> 70
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 Gly Pro Asp Ala Met Ile Leu Val Asp Gly Gln Pro Arg Gln Ala Lys
 20 25 30
 Gly Glu Leu
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 1 5 10 15
 Arg His Val

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Met Glu Met Ala Gly Gly Gly Pro Leu His Lys Phe Leu Val Gly Lys
1 5 10 15
Arg Glu Glu Ile
20

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Met Glu Phe Leu Pro Ser Gly Ser Leu Lys Glu Tyr Leu Pro Lys Asn
1 5 10 15

Lys Asn Lys Ile
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Lys Glu Arg Ile
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Met Glu Tyr Leu Pro Ser Gly Cys Leu Arg Asp Phe Leu Gln Arg His
1 5 10 15
Arg Ala Arg Leu
20

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Met Glu Tyr Val Pro Leu Gly Ser Leu Arg Asp Tyr Leu Pro Arg His
1 5 10 15
Ser Ile

<210> 77
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<400> 77

Leu	Glu	Tyr	Ala	Pro	Leu	Gly	Thr	Val	Tyr	Arg	Glu	Leu	Gln	Lys	Leu
1				5					10					15	

Ser Lys Phe

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Leu	Glu	Tyr	Cys	Ser	Gly	Gly	Glu	Leu	Phe	Asp	Arg	Ile	Glu	Pro	Asp
1				5					10					15	

Ile Gly Met

<210> 79

<211> 22

<212> PRT

<213> unknown

<220>

<223> IKK-1

<400> 79

Met	Glu	Tyr	Cys	Ser	Gly	Gly	Asp	Leu	Arg	Lys	Leu	Leu	Asn	Lys	Pro
1				5				10						15	

Glu Asn Cys Cys Gly Leu

20

<210> 80

<211> 22

<212> PRT

<213> unknown

<220>

<223> IKK-2

<400> 80

Met	Glu	Tyr	Cys	Gln	Gly	Gly	Asp	Leu	Arg	Lys	Tyr	Leu	Asn	Gln	Phe
1				5				10						15	

Glu Asn Cys Cys Gly Leu

20

<210> 81

<211> 19

<212> PRT
 <213> unknown

<220>
 <223> DAPK

<400> 81
 Leu Glu Leu Val Ala Gly Gly Glu Leu Phe Asp Phe Leu Ala Glu Lys
 1 5 10 15
 Glu Ser Leu

<210> 82
 <211> 31
 <212> PRT
 <213> unknown

<220>
 <223> IRK

<400> 82
 Met Glu Leu Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu
 1 5 10 15
 Arg Pro Glu Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
 20 25 30

<210> 83
 <211> 18
 <212> PRT
 <213> unknown

<220>
 <223> TGFbRII

<400> 83
 Thr Ala Phe His Ala Lys Gly Asn Leu Gln Glu Tyr Leu Thr Arg His
 1 5 10 15
 Val Ile

<210> 84
 <211> 18
 <212> PRT
 <213> unknown

<220>
 <223> ACTRIIA

<400> 84
 Thr Ala Phe His Glu Lys Gly Ser Leu Ser Asp Phe Leu Lys Ala Asn
 1 5 10 15
 Val Val

<210> 85
 <211> 18
 <212> PRT
 <213> unknown

<220>
 <223> ACTRIIB

<400> 85
 Thr Ala Phe His Asp Lys Gly Ser Leu Thr Asp Tyr Leu Lys Gly Asn
 1 5 10 15
 Ile Ile

210 86
 211 18
 212 PRT
 213 unknown

220
 223 ALK1

400 86
 Thr His Tyr His Glu His Gly Ser Leu Tyr Asp Phe Leu Gln Arg Gln
 1 5 10 15
 Thr Leu

210 87
 211 18
 212 PRT
 213 unknown

220
 223 ALK2

400 87
 Thr His Tyr His Glu Met Gly Ser Leu Tyr Asp Tyr Leu Gln Leu Thr
 1 5 10 15
 Thr Leu

210 88
 211 18
 212 PRT
 213 unknown

220
 223 ALK3

400 88
 Thr Asp Tyr His Glu Asn Gly Ser Leu Tyr Asp Phe Leu Lys Cys Ala

1
Thr Leu

5

10

15

<210> 89
<211> 18
<212> PRT
<213> unknown

<220>
<223> ALK4

<400> 89
Ser Asp Tyr His Glu His Gly Ser Leu Phe Asp Tyr Leu Asn Arg Tyr
1 5 10 15

Thr Val

CD

<400> 92
Thr Glu Tyr Met Glu Asn Gly Asp Leu Asn Gln Phe Leu Ser Arg His
1 5 10 15
Glu Pro

<210> 93
<211> 21
<212> PRT
<213> unknown

<220>
<223> ILK

<400> 93

Thr His Trp Met Pro Tyr Gly Ser Leu Tyr Asn Val Leu His Glu Gly
1 5 10 15
Thr Asn Phe Val Val
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<210> 94
<211> 16
<212> PRT
<213> unknown

<220>
<223> JNK

<400> 94
Met Glu Leu Met Asp Ala Asn Leu Cys Gln Val Ile Gln Met Glu Leu
1 5 10 15

<210> 95
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)
<223>

<221> AMIDATION
<222> (0)...(20)

<223> Akt1/Raca

<400> 95
Gly Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser Arg
1 5 10 15
Glu Arg Val Phe
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<210> 96
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(19)

<223> Alk1

<400> 96
Gly Thr His Tyr His Glu His Gly Ser Leu Tyr Asp Phe Leu Gln Arg
1 5 10 15

Gln Thr Leu

<210> 97
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(22)

<223> Braf

<400> 97
 Lys Lys Lys Lys Lys Lys Gly Ser Ser Leu Tyr His His Leu His
 1 5 10 15
 Ile Ile Glu Thr Lys Phe
 20

<210> 98
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)

<223> Braf

<400> 98
 Gly Thr Gln Trp Ser Glu Gly Ser Ser Leu Tyr His His Leu His Ile
 1 5 10 15
 Ile Glu Thr Lys Phe
 20

<210> 99
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(22)

<223> c-Ab1

<400> 99
 Gly Thr Glu Phe Met Thr Tyr Gly Asn Leu Leu Asp Tyr Leu Arg Glu
 1 5 10 15
 Cys Asn Arg Gln Glu Val
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<210> 100
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
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 <222> (1)...(0)

~~H~~ <221> AMIDATION
~~G~~ <222> (0)...(21)
~~G~~ <223>

~~G~~
~~G~~ <223> c-Met

<400> 100
~~G~~ Gly Leu Pro Tyr Met Lys His Gly Asp Leu Arg Asn Phe Ile Arg Asn
 1 5 10 15
~~G~~ Glu Thr His Asn Pro
 20

~~G~~
~~G~~ <210> 101
~~G~~ <211> 21
~~G~~ <212> PRT
 <213> Artificial Sequence

<220>
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<221> AMIDATION
 <222> (0)...(21)

<223> c-Raf

<400> 101
 Gly Thr Gln Trp Ser Glu Gly Ser Ser Leu Tyr Lys His Leu His Val
 1 5 10 15
 Gln Glu Thr Lys Phe
 20

<210> 102
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)

<223> benzyl ester at position 11

<221> AMIDATION

<222> (0)...(14)

<223> c-Raf

<400> 102

Ser Ser Leu Tyr Lys His Leu His Val Gln Glu Thr Lys Phe
 1 5 10

<210> 103

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(21)

<223> c-Sea

<400> 103

Gly Leu Pro Tyr Met Arg His Gly Asp Leu Arg His Phe Ile Arg Ala
 1 5 10 15
 Gln Glu Arg Ser Pro
 20

<210> 104

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> c-Src

<400> 104

Gly Thr Glu Tyr Met Ser Lys Gly Ser Leu Leu Asp Phe Leu Lys Gly
 1 5 10 15
 Glu Thr Gly Lys Tyr Leu
 20

<210> 105

<211> 14

<212> PRT

<213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 5
 benzyl ester at position 9

<221> AMIDATION
 <222> (0)...(14)

<223> c-Src

<400> 105
 Gly Ser Leu Leu Asp Leu Lys Gly Glu Thr Gly Lys Phe Leu
 1 5 10

<210> 106
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)
 <223>

<223> CDK2

<400> 106
 Gly Phe Glu Phe Leu His Gln Asp Leu Lys Lys Phe Met Asp Ala Ser
 1 5 10 15
 Ala Leu Thr Gly Ile
 20

<210> 107
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 1
 benzyl ester at position 7

<221> AMIDATION
 <222> (0)...(14)
 <223>

<223> CDK2

<400> 107

Asp Leu Lys Lys Phe Met Asp Ala Ser Ala Leu Thr Gly Met
1 5 10

<210> 108
<211> 14
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1) ... (0)
<223> benzyl ester at position 1
benzyl ester at position 7

<221> AMIDATION
<222> (0) ... (14)

<223> CDK4

<400> 108

Asp Leu Arg Thr Tyr Leu Asp Lys Ala Pro Pro Pro Gly Leu
1 5 10

<210> 109
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1) ... (0)

<221> AMIDATION
<222> (0) ... (21)

<223> CDK4

<400> 109

Gly Phe Glu His Val Asp Gln Asp Leu Arg Thr Tyr Leu Asp Lys Ala
1 5 10 15
Pro Pro Pro Gly Leu
20

<210> 110
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1) ... (0)

<221> AMIDATION
<222> (0) ... (21)

<223> CDK6

<400> 110
Gly Phe Glu His Val Asp Gln Asp Leu Thr Thr Tyr Leu Asp Lys Val
1 5 10 15
Pro Glu Pro Gly Val
20

<210> 111
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(19)

<223> Chk1

<400> 111
Gly Glu Tyr Ser Ser Gly Gly Glu Leu Phe Asp Arg Ile Glu Pro Asp
1 5 10 15
Ile Gly Met

<210> 112
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(19)
<223>

<223> Chk1

<400> 112
Gly Glu Tyr Ala Ser Gly Gly Glu Leu Phe Asp Arg Ile Glu Pro Asp
1 5 10 15
Ile Gly Met

<210> 113
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
 <222> (0)...(19)
 <223>

<223> CK IIa

<400> 113
 Lys Lys Lys Lys Lys Gly Gly Asn Asn Thr Asp Phe Lys Gln Leu Tyr
 1 5 10 15
 Gln Thr Leu

<210> 114
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(17)

<223> CK IIa

<400> 114
 Gly Phe Glu His Val Asn Asn Thr Asp Phe Lys Gln Leu Tyr Gln Thr
 1 5 10 15
 Leu

<210> 115
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(22)
 <223>

<223> Csk

<400> 115
 Gly Thr Glu Tyr Met Ala Lys Gly Ser Leu Val Asp Tyr Leu Arg Ser
 1 5 10 15
 Arg Gly Arg Ser Val Leu
 20

<210> 116

<211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 5

<221> AMIDATION
 <222> (0)...(14)

<223> Csk

<400> 116
 Gly Ser Leu Val Asp Leu Arg Ser Arg Gly Arg Ser Val Leu
 1 5 10

<210> 117
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)
 <221> AMIDATION
 <222> (0)...(21)

<223> Fak

<400> 117
 Gly Met Glu Leu Ser Thr Leu Gly Glu Leu Arg Ser Phe Leu Gln Val
 1 5 10 15
 Arg Lys Tyr Ser Leu
 20

<210> 118
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(17)

<223> FGFR-3

<400> 118
 Gly Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu
 1 5 10 15
 Glu

<210> 119
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1) ... (0)
 <223> benzyl ester at position 5
 benzyl ester at position 16

<221> AMIDATION
 <222> (0) ... (16)

<223> FGFR-3

<400> 119
 Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu Glu
 1 5 10 15

<210> 120
 <211> 23
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1) ... (0)
 <221> AMIDATION
 <222> (0) ... (23)

<223> FGFR-3

<400> 120
 Gly Val Glu Tyr Ala Ala Lys Gly Asn Leu Arg Glu Phe Leu Arg Ala
 1 5 10 15
 Arg Arg Pro Pro Gly Leu Glu
 20

<210> 121
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> stearyl at position 1

<221> AMIDATION
 <222> (0) ... (13)
 <223> FGFR-3

<400> 121
 Gly Ser Phe Asp Thr Ser Lys Pro Pro Glu Glu Gln Leu
 1 5 10

<210> 122
 <211> 23
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(23)

<223> Flk1

<400> 122
 Gly Val Glu Phe Ser Lys Phe Gly Asn Leu Ser Asn Phe Leu Arg Ala
 1 5 10 15
 Lys Arg Asn Leu Phe Val Pro
 20

<210> 123
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)
 <221> AMIDATION
 <222> (0)...(17)
 <223>

<223> Flk1

<400> 123
 Gly Gly Asn Leu Ser Asn Phe Leu Arg Ala Lys Arg Asn Leu Phe Val
 1 5 10 15
 Pro

<210> 124
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(16)

<223> Flk1

<400> 124
Gly Asn Leu Ser Asn Phe Leu Arg Ala Lys Arg Asn Leu Phe Val Pro
1 5 10 15

<210> 125
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> stearyl at position 1

<221> AMIDATION
<222> (0)...(13)
<223> Flk1

<400> 125
Gly Arg Phe Arg Gln Gly Lys Asp Tyr Val Gly Glu Leu
1 5 10

<210> 126
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<221> AMIDATION
<222> (0)...(22)
<223> GSK3b

<400> 126
Lys Lys Lys Lys Lys Gly Gly Val Ala Arg His Tyr Ser Arg
1 5 10 15
Ala Lys Gln Thr Leu Pro
20

<210> 127
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<221> AMIDATION
<222> (0)...(13)
<223> GSK3b

<400> 127
Val Ala Arg His Tyr Ser Arg Ala Lys Gln Thr Leu Pro
1 5 10

<210> 128
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> GSK3b

<400> 128

Gly Asp Tyr Val Pro Glu Thr Val Tyr Arg Val Ala Arg His Tyr Ser
1 5 10 15
Arg Ala Lys Gln Thr Leu
20

<210> 129
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(12)

<223> GSK3b

<400> 129
Arg Val Ala Arg His Tyr Ser Arg Ala Lys Gln Thr
1 5 10

<210> 130
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> Hck

<400> 130
Gly Thr Glu Phe Met Ala Lys Gly Ser Leu Leu Asp Phe Leu Lys Ser
1 5 10 15
Asp Glu Gly Ser Lys Gln
20

<210> 131
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION

<222> (0) . . . (20)

<223> Tak1

<400> 131
Gly Leu Glu Tyr Ala Pro Leu Gly Thr Val Tyr Arg Glu Leu Gln Lys
1 5 10 15
Leu Ser Lys Phe
20

<210> 132

<211> 23

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1) . . . (0)

<221> AMIDATION

<222> (0) . . . (23

1222 TIAN

<400> 132

Gly Met Glu Tyr Ser Ser Gly Gly Asp Leu Arg Lys Leu Leu Asn Lys
1 5 10 15
Pro Glu Asn Ser Ser Gly Leu
20

<210> 133

<211> 23

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1) . . . (0)

<221> AMIDATION

<222> (0) . . . (23)

<223>

<223> IKK-2

<400> 133

Gly Met Glu Tyr Ser Gln Gly Gly Asp Leu Arg Lys Tyr Leu Asn Gln
 1 5 10 15
 Phe Glu Asn Ser Ser Gly Leu
 20

<210> 134
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
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<221> AMIDATION
 <222> (0)...(22)

<223> ILK

<400> 134
 Gly Thr His Trp Met Pro Tyr Gly Ser Leu Tyr Asn Val Leu His Glu
 1 5 10 15
 Gly Thr Asn Phe Val Val
 20

<210> 135
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> stearyl at position 1

<221> AMIDATION
 <222> (0)...(13)
 <223> ILK

<400> 135
 Gly Tyr Asn Val Leu His Glu Gly Thr Asn Phe Val Val
 1 5 10

<210> 136
 <211> 19
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(19)
 <223>

<223> IRK

<400> 136
Gly Met Glu Leu Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser
1 5 10 15
Leu Arg Pro

<210> 137
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(12)

<223> IRK

<400> 137
Ala Gln Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
1 5 10

<210> 138
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> IRK

<400> 138
Gly Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu Ala
1 5 10

<210> 139
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
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<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> IRK

<400> 139
Gly Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
1 5 10

<210> 140
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(17)

<223> IRK

<400> 140
Gly Leu Arg Pro Glu Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr
1 5 10 15
Leu

<210> 141
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(21)

<223> Jak1

<400> 141
Gly Met Glu Phe Leu Pro Ser Gly Ser Leu Lys Glu Tyr Leu Pro Lys
1 5 10 15
Asn Lys Asn Lys Ile
20

<210> 142
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> Jak1

<400> 142
Gly Leu Lys Glu Tyr Leu Pro Lys Asn Lys Asn Lys Ile
1 5 10

<210> 143
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> Jak2

<400> 143
Gly Leu Arg Asp Tyr Leu Gln Lys His Lys Glu Arg Ile
1 5 10

<210> 144
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> stearyl at position 1

<221> AMIDATION
<222> (0)...(11)
<223> Jak2

<400> 144
Gly Leu Arg Asp Tyr Leu Gln Lys His Lys Glu
1 5 10

<210> 145
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(20)

<223> Jak3

<400> 145

Gly Met Glu Tyr Leu Pro Ser Gly Ser Leu Arg Asp Phe Leu Gln Arg
1 5 10 15
His Arg Ala Leu
20

<210> 146

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(21)

<223> Jak3

<400> 146

Gly Met Glu Tyr Leu Pro Ser Gly Ser Leu Arg Asp Phe Leu Gln Arg
1 5 10 15
His Arg Ala Arg Leu
20

<210> 147

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(13)

<223> Jak3

<400> 147

Gly Leu Arg Asp Phe Leu Gln Arg His Arg Ala Arg Leu
1 5 10

<210> 148

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<221> ACETYLATION

<222> (1)...(0)
<223> benzyl ester at position 5

<221> AMIDATION
<222> (0)...(14)

<223> Lck

<400> 148
Gly Ser Leu Val Asp Leu Lys Thr Pro Ser Gly Ile Lys Leu
1 5 10

<210> 149
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> Lck

<400> 149
Gly Thr Glu Tyr Met Glu Asn Gly Ser Leu Val Asp Phe Leu Lys Thr
1 5 10 15
Pro Ser Gly Ile Lys Leu
20

<210> 150
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> Lyn

<400> 150
Gly Thr Glu Tyr Met Ala Lys Gly Ser Leu Leu Asp Phe Leu Lys Ser
1 5 10 15
Asp Glu Gly Gly Lys Val
20

<210> 151
<211> 20
<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(20)

<223> MARK1

<400> 151

Gly Met Glu Tyr Ala Ser Gly Gly Glu Val Phe Asp Tyr Leu Val Ala
 1 5 10 15
 His Gly Arg Met
 20

<210> 152

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<221> ACETYLATION

<222> (1)...(0)

<223> benzyl ester at position 2

benzyl ester at position 5

<221> AMIDATION

<222> (0)...(15)

<223> PDGFR-b

<400> 152

Gly Asp Leu Val Asp Tyr Leu His Arg Asn Lys His Thr Phe Leu
 1 5 10 15

<210> 153

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> PDGFR-b

<400> 153

Gly Thr Glu Tyr Ser Arg Tyr Gly Asp Leu Val Asp Tyr Leu His Arg
 1 5 10 15
 Asn Lys His Thr Phe Leu
 20

<210> 154
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(20)

<223> PKC_b

<400> 154
Gly Met Glu Tyr Val Asn Gly Gly Asp Leu Met Tyr His Ile Gln Gln
1 5 10 15
Val Gly Arg Phe
20

<210> 155
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(20)

<223> PKC_b

<400> 155
Lys Lys Lys Lys Lys Gly Gly Asp Leu Met Tyr His Ile Gln Gln
1 5 10 15
Val Gly Arg Phe
20

<210> 156
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<223> benzyl ester at position 5

<221> AMIDATION
<222> (0)...(12)

<223> Plk

<400> 156
Arg Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala

1

5

10

<210> 157
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)
<223> benzyl ester at position 6

<221> AMIDATION
<222> (0)...(13)

<223> Plk

<400> 157
Gly Arg Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala
1 5 10

<210> 158
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)
<221> AMIDATION
<222> (0)...(20)

<223> Plk

<400> 158
Gly Leu Glu Leu Ser Arg Arg Arg Ser Leu Leu Glu Leu His Lys Arg
1 5 10 15
Arg Lys Ala Leu
20

<210> 159
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)
<221> AMIDATION
<222> (0)...(22)

<223> Ret

<400> 159

Gly Val Glu Tyr Ala Lys Tyr Gly Ser Leu Arg Gly Phe Leu Arg Glu
1 5 10 15
Ser Arg Lys Val Gly Pro
20

<210> 160
<211> 15
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)
<223> benzyl ester at position 9

<221> AMIDATION
<222> (0)...(15)

<223> Ret

<400> 160

Gly Ser Leu Arg Gly Phe Leu Arg Glu Ser Arg Lys Val Gly Pro
1 5 10 15

<210> 161
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(21)

<223> Ron

<400> 161

Gly Leu Pro Tyr Met Cys His Gly Asp Leu Leu Gln Phe Ile Arg Ser
1 5 10 15
Pro Gln Arg Asn Pro
20

<210> 162
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(20)

<223> SNK

<400> 162

Gly Leu Glu Tyr Ser Ser Arg Arg Ser Met Ala His Ile Leu Lys Ala
 1 5 10 15
 Arg Lys Val Leu
 20

<210> 163

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(20)

<223> Syk

<400> 163

Gly Met Glu Met Ala Glu Leu Gly Pro Leu Asn Lys Tyr Leu Gln Gln
 1 5 10 15
 Asn Arg His Val
 20

<210> 164

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(19)

<223> TGFbRII

<400> 164

Gly Thr Ala Phe His Ala Lys Gly Asn Leu Gln Glu Tyr Leu Thr Arg
 1 5 10 15
 His Val Ile

<210> 165

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION
<222> (0)...(25)

<223> TrkB

<400> 165
Gly Phe Glu Tyr Met Lys His Gly Asp Leu Asn Lys Phe Leu Arg Ala
1 5 10 15
His Gly Pro Asp Ala Val Leu Met Ala
20 25

<210> 166
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> TrkB

<400> 166
Gly Leu Arg Ala His Gly Pro Asp Ala Val Leu Met Ala
1 5 10

<210> 167
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(11)

<223> TrkB

<400> 167
Gly Leu Arg Ala His Gly Pro Asp Ala Val Leu
1 5 10

<210> 168
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> TrkB

<400> 168
Gly Leu Asn Phe Lys Leu Arg Ala His Gly Pro Asp Ala
1 5 10

<210> 169
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(13)

<223> TrkB

<400> 169
Gly Phe Lys Leu Arg Ala His Gly Pro Asp Ala Val Leu
1 5 10

<210> 170
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(21)

<223> Zap70

<400> 170
Gly Met Glu Met Ala Gly Gly Gly Pro Leu His Lys Phe Leu Val Gly
1 5 10 15
Lys Arg Glu Glu Ile
20

<210> 171
<211> 21
<212> PRT
<213> Unknown

<220>
<223> IRK

<400> 171
Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu
1 5 10 15
Ala Glu Asn Asn Pro
20

<210> 172
<211> 8
<212> PRT
<213> Unknown

<220>
<223> endothelial growth factor receptor

<400> 172
Lys Phe Asp Val Ile Asn Leu Ala
1 5